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(54) Title: DNA ENCODING THE HUMAN P2U RECEPTOR AND NULL CELLS EXPRESSING P2U RECEPTORS

(57) Abstract

Isolated DNAs encoding the human P2v receptor are disclosed, along with vectors and host cells containing the same and methods of using the same. Host cells which are essentially free of endogenous P2U receptor expression, and which express a heterologous P2U receptor such as a murine P2U receptor, are also disclosed, along with methods of using the same.

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DNA ENCODING THE HUMAN P_{2U} RECEPTOR AND NULL CELLS EXPRESSING P_{2U} RECEPTORS

Field of the Invention

The present invention relates to DNA encoding P_{2U} receptors, and particularly relates to DNA encoding the human P_{2U} receptor, and null cells which express heterologous P_{2U} receptors.

Background of the Invention

Cystic fibrosis (CF) exhibits reduced Cl secretion by airway epithelia. Consequently, one of the most debilitating effects of CF is the development of a dehydrated, viscous mucus which obstructs the airways and compromises lung function. Extracellular nucleotide triphosphates, such as ATP or UTP, are able to regulate Clsecretion in human airway epithelia and, combination with an inhibitor of Na* transport, provide an alternative, non-CFTR-dependent mechanism to fluid secretion in CF airway epithelia. Extracellular nucleotides also stimulate mucus secretion by goblet cells in vitro and excessive activation of this pathway in vivo may be partly responsible for hypersecretion observed in chronic bronchitis. In both 20 cases, the responses are mediated by 5'-nucleotide (P_{211}) receptors on the cell surface.

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Recently, a cDNA encoding a murine ATP/UTP receptor was cloned from neuroblastoma cells by functional expression in Xenopus oocytes (K. Lustig et al., Proc. Natl. Acad. Sci. USA 90, 5113 (1993)). The receptor, a member of the G protein-coupled receptor superfamily, is activated by UTP and ATP, initiates elevation of cytoplasmic calcium, and has been identified with the subtype of P2-receptor provisionally designated P2U. Its pharmacological and signaling properties are very similar to those described for a 5'-nucleotide (P2U) receptor present in the human airway epithelial cell line, CF/T43, which was derived from a donor with CF (A. Brown et al., Mol. Pharmacol. 40, 648 (1991)).

Isolation and molecular characterization of the receptor for extracellular nucleotides present in human airway epithelia will permit studies of the expression of this receptor in normal and diseased tissues and facilitate identification of new drugs for therapy.

Summary of the Invention

In view of the foregoing, a first aspect of the present invention is isolated DNA encoding an P_{2U} receptor selected from the group consisting of: (a) isolated DNA (e.g., the DNA of SEQ ID NO:1) which encodes the human P_{2U} receptor having the amino acid sequence given herein as SEQ ID NO:2; (b) isolated DNA which hybridizes to isolated DNA of (a) above and which encodes a human P_{2U} receptor; and (c) isolated DNA differing from the isolated DNAs of (a) and (b) above in nucleotide sequence due to the degeneracy of the genetic code, and which encodes a human P_{2U} receptor.

A second aspect of the present invention is a recombinant DNA sequence comprising vector DNA and a DNA as given above which encodes an P_{20} receptor.

A third aspect of the present invention is a host cell containing a recombinant DNA sequence as given above and capable of expressing the encoded P_{20} receptor.

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A fourth aspect of the present invention is isolated DNA as given above, and oligonucleotides as given above, configured in antisense for the production of antisense RNA which inhibits the expression of P_{2U} 5 receptor. (hereinafter referred to as "antisense DNAs"). Such antisense DNAs may be provided in a vector as given herein for transcription in a suitable cell where they then inhibit the production of the P_{20} receptor. alternative, antisense oligonucleotides which bind to RNA in cells and inhibit the expression of $P_{2\text{U}}$ receptor therein may be delivered directly to cells.

A fifth aspect of the present invention is a transformed null cell which is essentially free of endogenous P_{2u} receptor expression, which null cell contains and expresses heterogenous DNA encoding an P_{2U} receptor, said heterogenous DNA selected from the group consisting of: (a) isolated DNA consisting essentially of DNA which encodes the human P20 receptor having the amino acid sequence given herein as SEQ ID NO:2; (b) isolated DNA which hybridizes to isolated DNA of (a) above and 20 which encodes a P_{2U} receptor; and (c) isolated DNA differing from the isolated DNAs of (a) and (b) above in nucleotide sequence due to the degeneracy of the genetic code, and which encodes a Pzu receptor.

25 A sixth aspect of the present invention is a method of detecting compounds which bind to the P_{2U} The method comprises: (a) providing a cell receptor. which contains and expresses heterogenous DNA encoding a heterogenous P_{2U} receptor, which cell secretes endogenous ATP in response to stimulation of said P_{20} receptor, the 30 heterogenous DNA being as described above; (b) removing essentially all endogenous ATP secreted by said cell from said cell; (c) contacting a compound to said cell; and then (d) detecting the binding of the compound to the P_{2U} 35 receptor (e.g., by detecting an elevation intracellular calcium therein).

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A seventh aspect of the present invention is a method of enhancing cell growth, comprising transforming a cell with a vector capable of expressing a P2U receptor in the cell in an amount sufficient to enhance cell growth by autostimulation of the P20 receptor, where the cell secretes ATP which stimulates the P_{2U} receptor, and where cell growth is enhanced by stimulation of the P2U receptor. The method is useful, among other things, for stimulating the growth of cultures of mammalian cells in vitro where the cells are difficult to grow or otherwise grow slowly in culture.

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An eighth aspect of the present invention is a method of inhibiting cell growth, comprising transforming a cell with a vector capable of expressing a P_{2U} receptor in the cell in an amount sufficient to inhibit cell growth by autostimulation of the P_{2U} receptor, where the cell secretes ATP, and where cell growth is inhibited by stimulation of the P_{2U} receptor. The method is useful, among other things, for inhibiting the growth of cells in 20 vitro or in vivo such as in proliferative or hyperplastic diseases, both malignant and nonmalignant, of human and animal subjects.

A ninth aspect of the present invention is a method of treating diseases of epithelial surfaces (particularly diseases of airway epithelial mucosal surfaces such as cystic fibrosis, asthma, and chronic bronchitis) in a human or animal subject in need of such treatment by enhancing or inhibiting mucus production in that subject. The method is carried out by administering to the subject a vector capable of entering lung epithelial cells. The vector may be either:

> (a) a vector which carries a DNA encoding a P_{2U} receptor as described herein, which DNA is operably associated with a promoter which expresses that gene in the epithelial cells; or

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(b) a vector containing an antisense nucleic acid which encodes an antisense RNA, which antisense RNA is capable of binding to pre-mRNA encoding a P_{2U} receptor and inhibiting expression thereof in the epithelial cells.

Also disclosed herein is a vector as described above, pharmaceutical formulations containing such a vector, and the use of such vectors for the preparation of a medicament the treatment of epithelial mucosal diseases in a subject in need of such treatment.

The foregoing and other aspects of the present invention are explained in detail below.

Brief Description of the Drawings

Figure 1A shows [3H]inositol phosphate

15 production in cpm x 10⁻³ in wild type, LSN, and LHP2USN cells, with and without the present of apyrase in cells which had not been exposed to exogenous nucleotides.

Figure 1B shows [3 H]inositol phosphate production in cpm x 10^{-3} in wild type, LSN, and LHP2USN cells, in cells which had been exposed to exogenous carbachol, UTP, and ATP, or cells which had not been exposed to exogenous compounds (control).

Figure 2 shows the effect of nucleotides and nucleotide analogs on $\{Ca^{2+}\}_i$ in K-562 cells stably expressing a P_{2U} receptor. $\{Ca^{2+}\}_i$ was measured after treatment with the indicated concentration of UTP (filled circles), ATP (open circles), ATP γ S (filled triangles), BzATP (open triangles), ATP α S (filled squares), or 2-MeSATP (open squares).

30 Detailed Description of the Invention

Amino acid sequences disclosed herein are presented in the amino to carboxy direction, from left to right. The amino and carboxy groups are not presented in the sequence. Nucleotide sequences are presented herein by single strand only, in the 5' to 3' direction, from

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left to right. Nucleotides and amino acids are represented herein in the manner recommended by the IUPAC-IUB Biochemical Nomenclature Commission, or (for amino acids) by three letter code, in accordance with 37 CFR §1.822 and established usage. See, e.g., PatentIn User Manual, 99-102 (Nov. 1990) (U.S. Patent and Trademark Office, Office of the Assistant Commissioner for Patents, Washington, D.C. 20231); U.S. Patent No. 4,871,670 to Hudson et al. at Col. 3 lines 20-43 (applicants specifically intend that the disclosure of this and all other patent references cited herein be incorporated herein by reference).

Subjects treated by methods disclosed herein include both human subjects and animal subjects (e.g., dog, cat, horse) for veterinary purposes.

A. DNAs ENCODING P_{2U} RECEPTORS AND OLIGONUCLEOTIDES THEREOF

DNAs which encode P_{2U} receptors, whether they are cDNAs or genomic DNAs, encode a G-protein coupled receptor protein which, on expression in a suitable host cell, (a) selectively and stereospecifically binds ATP and UTP and (b) initiates the elevation of cytoplasmic calcium on binding ATP or UTP. This definition is intended to encompass natural allelic variations in the DNAs.

DNAs encoding P_{2U} receptors which hybridize to the DNA encoding the human P_{2U} receptor disclosed herein, may be of any species of origin, including murine (mouse, rat), rabbit, cat, porcine, and human, but preferably code for an P_{2U} receptor of mammalian origin, and most preferably code for human P_{2U} receptors.

Hybridization conditions which will permit other DNA sequences which code on expression for an P_{2U} receptor to hybridize to a DNA sequence as given herein are, in general, high stringency conditions. For example, hybridization of such sequences may be carried out under conditions represented by a wash stringency of

0.3 M NaCl, 0.03 M sodium citrate, 0.1% SDS at 60°C or even 70°C to DNA disclosed herein in a standard in situ hybridization assay. (See J. Sambrook et al., Molecular Cloning, A Laboratory Manual (2d Ed. 1989) (Cold Spring Harbor Laboratory)). The same hybridization conditions are used to determine hybridization of oligonucleotides.

In general, DNA sequences which code for P_{2U} receptors and hybridize to the DNA sequence encoding the human P_{2U} receptor disclosed herein will be at least 70%, 10 75%, 80%, 85%, 90%, or even 95% homologous or more with the sequence of the DNA encoding the human P_{2U} receptor disclosed herein.

In general, DNA sequences which encode human P_{2U} receptors which hybridize to the DNA encoding the human 15 P_{2U} receptor disclosed herein will be 93%, 94%, 95%, or even 97% homologous or more to the DNA sequence encoding the human P_{2U} receptor disclosed herein.

Further, DNA sequences which code for the same P_{2U} receptor as coded for by the foregoing sequences, but which differ in codon sequence from these due to the degeneracy of the genetic code, are also an aspect of this invention. The degeneracy of the genetic code, which allows different nucleic acid sequences to code for the same protein or peptide, is well known in the literature. See e.g., U.S. Patent No. 4,757,006 to Toole et al. at Col. 2, Table 1.

B. GENETIC ENGINEERING TECHNIQUES

The production of cloned genes, recombinant DNA, vectors, transformed host cells, proteins and protein fragments by genetic engineering is well known. See, e.g., U.S. Patent No. 4,761,371 to Bell et al. at Col. 6 line 3 to Col. 9 line 65; U.S. Patent No. 4,877,729 to Clark et al. at Col. 4 line 38 to Col. 7 line 6; U.S. Patent No. 4,912,038 to Schilling at Col. 3 line 26 to Col. 14 line 12; and U.S. Patent No. 4,879,224 to Wallner at Col. 6 line 8 to Col. 8 line 59.

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A vector is a replicable DNA construct. Vectors are used herein either to amplify DNA encoding P20 receptors as given herein and/or to express DNA which encodes P_{2U} receptors as given herein. An expression 5 vector is a replicable DNA construct in which a DNA sequence encoding a P2U receptor is operably linked to suitable control sequences capable of effecting the expression of the receptor in a suitable host. The need for such control sequences will vary depending upon the host selected and the transformation method chosen. Generally, control sequences include a transcriptional promoter, an optional operator sequence to control sequence encoding suitable transcription, a mRNA ribosomal binding sites, and sequences which control the termination of transcription and translation.

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Amplification vectors do not require expression control domains. All that is needed is the ability to replicate in a host, usually conferred by an origin of replication, and a selection gene to facilitate recognition of transformants.

Vectors comprise plasmids (e.g., the neomycinresistance plasmid pP2R1 for stable transfection; the plasmids pCCM6x1 or pP2R for transient expression), viruses (e.g., adenovirus, cytomegalovirus), phage, and integratable DNA fragments (i.e., fragments integratable into the host genome by recombination). The vector replicates and functions independently of the host genome, or may, in some instances, integrate into the genome itself. Expression vectors should contain a promoter and RNA binding sites which are operably linked to the gene to be expressed and are operable in the host organism.

DNA regions are operably linked or operably associated when they are functionally related to each other. For example, a promoter is operably linked to a coding sequence if it controls the transcription of the sequence; or a ribosome binding site is operably linked

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to a coding sequence if it is positioned so as to permit translation.

Transformed host cells are cells which have been transformed or transfected with vectors containing 5 a DNA sequence as disclosed herein constructed using recombinant DNA techniques. Transformed host cells ordinarily express the receptor, but host cells transformed for purposes of cloning or amplifying the receptor DNA do not need to express the receptor.

Suitable host cells include prokaryote, yeast 10 or higher eukaryotic cells such as mammalian cells and insect cells. Cells derived from multicellular organisms are a particularly suitable host for recombinant P20 receptor synthesis, and mammalian cells are particularly preferred. Propagation of such cells in cell culture has 15 become a routine procedure (Tissue Culture, Academic Press, Kruse and Patterson, editors (1973)). Examples of useful host cell lines are VERO and HeLa cells, Chinese hamster ovary (CHO) cell lines, and WIL38, BHK, COS-7, 20 CV, and MDCK cell lines. Expression vectors for such cells ordinarily include (if necessary) an origin of replication, a promoter located upstream from the DNA encoding the P_{2U} receptor to be expressed and operatively associated therewith, along with a ribosome binding site, 25 an RNA splice site (if intron-containing genomic DNA is used), a polyadenylation site, and a transcriptional termination sequence.

The transcriptional and translational control sequences in expression vectors to be used in transforming vertebrate cells are often provided by viral sources. For example, commonly used promoters are derived from polyoma, Adenovirus 2, and Simian Virus 40 (SV40). See, e.g., U.S. Patent No. 4,599,308.

An origin of replication may be provided either 35 by construction of the vector to include an exogenous origin, such as may be derived from SV40 or other viral source (e.g. Polyoma, Adenovirus, VSV, or BPV), or may be

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provided by the host cell chromosomal replication mechanism. If the vector is integrated into the host cell chromosome, the latter is often sufficient. Rather than using vectors which contain viral origins of replication, one can transform mammalian cells by the method of cotransformation with a selectable marker and the receptor DNA. Examples of suitable selectable markers are dihydrofolate reductase (DHFR) or thymidine kinase. This method is further described in U.S. Pat.

10 No. 4,399,216.

Other methods suitable for adaptation to the synthesis of the P_{2U} receptor in recombinant vertebrate cell culture include those described in M-J. Gething et al., Nature 293, 620 (1981); N. Mantei et al., Nature 281, 40; A. Levinson et al., EPO Application Nos. 117,060A and 117,058A.

Host cells such as insect cells (e.g., cultured Spodoptera frugiperda cells) and expression vectors such as the baculovirus expression vector (e.g., vectors derived from Autographa californica MNPV, Trichoplusia ni MNPV, Rachiplusia ou MNPV, or Galleria ou MNPV) may be employed in carrying out the present invention, as described in U.S. Patents Nos. 4,745,051 and 4,879,236 to Smith et al. In general, a baculovirus expression vector comprises a baculovirus genome containing the gene to be expressed inserted into the polyhedrin gene at a position ranging from the polyhedrin transcriptional start signal to the ATG start site and under the transcriptional control of a baculovirus polyhedrin promoter.

Prokaryote host cells include gram negative or gram positive organisms, for example Escherichia coli (E. coli) or Bacilli. Higher eukaryotic cells include established cell lines of mammalian origin as described below. Exemplary host cells are E. coli W3110 (ATCC 27,325), E. coli B, E. coli X1776 (ATCC 31,537), E. coli 294 (ATCC 31,446). A broad variety of suitable prokaryotic and microbial vectors are available. E. coli

is typically transformed using pBR322. Promoters most commonly used in recombinant microbial expression vectors include the beta-lactamase (penicillinase) and lactose promoter systems (Chang et al., Nature 275, 615 (1978); and Goeddel et al., Nature 281, 544 (1979)), a tryptophan (trp) promoter system (Goeddel et al., Nucleic Acids Res. 8, 4057 (1980) and EPO App. Publ. No. 36,776) and the tac promoter (H. De Boer et al., Proc. Natl. Acad. Sci. USA 80, 21 (1983)). The promoter and Shine-Dalgarno sequence (for prokaryotic host expression) are operably linked to the DNA encoding the P2U receptor, i.e., they are positioned so as to promote transcription of P2U receptor messenger RNA from the DNA.

Eukaryotic microbes such as yeast cultures may also be transformed with vectors carrying the isolated 15 DNA's disclosed herein. see, e.g., U.S. Patent No. 4,745,057. Saccharomyces cerevisiae is the most commonly used among lower eukaryotic host microorganisms, although a number of other strains are commonly available. 20 vectors may contain an origin of replication from the 2 micron yeast plasmid or an autonomously replicating sequence (ARS), a promoter, DNA encoding the receptor as given herein, sequences for polyadenylation transcription termination, and a selection gene. exemplary plasmid is YRp7, (Stinchcomb et al., Nature 282, 39 (1979); Kingsman et al., Gene 7, 141 (1979); Tschemper et al., <u>Gene</u> <u>10</u>, 157 (1980)). promoting sequences in yeast vectors include promoters for metallothionein, 3-phosphoglycerate kinase (Hitzeman et al., <u>J. Biol. Chem.</u> 255, 2073 (1980) or 30 other glycolytic enzymes (Hess et al., <u>J. Adv. Enzyme</u> Req. 7, 149 (1968); and Holland et al., Biochemistry 17, 4900 (1978)). Suitable vectors and promoters for use in yeast expression are further described in R. Hitzeman et al., EPO Publn. No. 73,657. 35

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C. USES OF DNAS ENCODING P2U RECEPTORS.

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P₂₀ receptors made from cloned genes accordance with the present invention may be used for screening compounds for P_{2U} receptor activity, or for determining the amount of a P2U receptor agonist antagonist such as ATP or UTP in a solution (e.g., blood for or serum), either the purpose quantitatively measuring the presence of the agonist or in the solution (e.g.,antagonist in a laboratory assay), or screening for the presence or absence of that compound in the solution. For example, host cells may be transformed with a vector of the present invention, P20 receptors expressed in that host, the cells lysed, and the membranes from those cells used to screen compounds for P_{2U} receptor binding activity. Competitive binding assays in which such procedures may be carried out are well known.

By selection of "null" host cells which do not ordinarily express P2U receptors (e.g., cells which are essentially free of endogenous P_{2U} receptor expression, and more preferably cells which are essentially free of endogenous P_{2x} , P_{2y} and P_{2u} receptor expression, which cells may or may not be essentially free of endogenous P2T receptor expression), preparations free of other P2 receptors which might interfere with the assay can be obtained. Further, Pau receptor agonists and antagonists, particularly UTP and analogs thereof, can be identified by transforming host cells with vectors of the present invention, which host cells also express the several Gprotein subunits and effector proteins necessary to cause the cell to initiate elevation of cytoplasmic calcium. agonists will initiate elevation receptor cytoplasmic calcium when they bind to and activate receptors expressed in such cells. Such cells must be capable of operatively associating the P2U receptor with the appropriate effector proteins: e.g., G protein must also be present in the cell membranes in the appropriate

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configuration. Such cells are typically mammalian cells, including human cells, examples being human leukemia cells (e.g., K562 human leukemia cells), and human astrocytoma cells (e.g., 1321N, human astrocytoma cells).

As noted above, one particular method of detecting compounds which bind to the P_{2U} receptor comprises (a) providing a cell which contains expresses heterogenous DNA encoding a heterogenous Pau receptor, which cell secretes endogenous ATP which stimulates the P_{2U} receptor, the heterogenous DNA being as described above; (b) removing essentially all endogenous ATP secreted by the cell from the cell (e.g., inactivating the endogenous ATP or by separating the endogenous ATP from the cell); (c) contacting a compound to the cell; and then (d) detecting the binding of the compound to the P2U receptor. The method is particularly useful for detecting UTP and analogs thereof. contacting step is typically carried out in an aqueous solution, though the cells can also be immobilized on a solid support for the assay. The removing step may be carried out by any suitable means, such as by including an enzyme in the solution which utilizes ATP as a substrate (e.g., including hexokinase and glucose in the aqueous solution in an amount sufficient so that essentially all of said ATP is utilized to phosphorylate the glucose; by including apyrase in the aqueous solution in an amount effective to degrade the ATP; etc.). Alternatively, the removing step may be carried out by washing the cell. The binding of the compound to the Pau receptor and activation of the receptor by agonists thereof may be detected by any suitable means, such as measuring for or detecting an elevation in intracellular calcium, measuring for or detecting an elevation in intracellular inositol phosphates, etc. Where only an indication of binding is desired, detection of binding may be carried out by any suitable means, such as with a

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competitive binding assay with a known agonist or antagonist.

Vectors which are capable of expressing a P211 receptor in a host cell are useful in methods of 5 enhancing, facilitating, and/or accelerating cell growth. Such methods involve the step of transforming a cell with a vector capable of expressing a P_{2U} receptor in the cell in an amount sufficient to enhance cell growth. growth is enhanced by autostimulation of the P_{2U} receptor, wherein the cell secretes ATP which activates the P_{2U} receptor, and wherein cell growth is enhanced activation of the P_{2U} receptor (i.e., the binding of an agonist thereto). The vector may be as described above. The method is useful, among other things, for stimulating the growth of cultures of mammalian cells in vitro where the cells are difficult to grow or otherwise grow slowly in culture.

Vectors which are capable of expressing a P20 receptor in a host cell are also useful in methods of inhibiting or slowing cell growth. Such methods involve the step of transforming a cell with a vector capable of expressing a P₂₀ receptor in the cell in an amount sufficient to inhibit or slow cell growth. Cell growth is inhibited by autostimulation of the P_{20} receptor, wherein the cell secretes ATP, and wherein cell growth is inhibited by activation of the P_{2U} receptor. The vector may again be as described above. The method is useful, among other things, for slowing the growth cells in vitro where the cells are unduly proliferative, and for combatting proliferative or hyperplastic growth of cells or cell populations in a subject in need of such treatment, including both malignant and non-malignant cell growth (e.g., goblet cell hyperplasia).

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Another aspect of the invention is antisense oligonucleotides (and DNAs encoding the same) having a sequence capable of binding specifically with any sequences of an mRNA molecule which encodes a human P_{2U}

receptor so as to prevent translation of the mRNA molecule (binding conditions may be at the stringencies as given above with respect to DNA hybridization). method may be carried out in any patient where decreasing mucus secretion by downregulating epithelial cell (e.g., lung airway epithelial cell) cell P_{2U} receptor expression is desired. This method may also be carried out in vitro or in a subject in need of such treatment for the purpose of slowing cell growth for the purposes described above. For example, in chronic bronchitis, the airway cell 10 distribution is changed so that the number of goblet cells is increased and the number of ciliated cells is This phenomenon, known as goblet cell decreased. hyperplasia, is caused by chronic toxic exposure, and in patients afflicted with this disorder a means for 15 decreasing mucus secretion by inhibiting P_{2U} receptor expression in airway epithelial cells by administration of antisense nucleotides as described herein is desired. Other examples where such treatment may be desired include the treatment of gallblader epithelial mucos for 20 cholecystitis and the treatment of liver biliary ducts Where the antisense olgionucleotides for cholangitis. are delivered directly to cells rather than via a DNA intermediate, nucleotides in which the phosphodiester bonds have been modified, e.g., to the methylphosphonate, 25 phosphotriester, the phosphorothioate, phosphorodithioate, or the phosphoramidate, so as render the oligonucleotide more stable in vivo may be used. Antisense oligonucleotides may be of any suitable length (e.g., from about 8 or 10 to 50 or 60 nucleotides 30 in length), depending on the particular target being bound and the mode of delivery thereof. The antisense oligonucleotides are, in general, delivered or dosaged so that intracellular levels of the antisense oligonucleotide of from .05 to 50 μM are achieved. 35

Antisense oligonucleotides may be combined with an appropriate physiologically or pharmaceutically

carrier (e.g., sterile acceptable pyrogen-free solution) saline to provide physiological pharmaceutical formulation for delivery of the antisense oligonucleotide to cells, also in accordance with known See, e.g., U.S. Patent No. 5,023,243 to techniques. antisense oligonucleotide The Tullis). formulation may be enclosed within a liposome, microcrystal, or other lipid vesicle to facilitate its being carried into the interior of cells, in accordance with known techniques, as also discussed 10 Antisense oligonucleotides may also be delivered to cells by way of a DNA intermediate, as discussed in greater detail below.

In one embodiment, the present invention is carried out by administering to the subject a vector carrying a nucleic acid active agent, which vector is capable of entering epithelial mucosa cells (e.g., lung airway epithelial mucosa cells). Such vectors may be formulated with pharmaceutical carriers and administered topically to the epithelial cells, such as by use of an 20 aerosol delivery system in which respirable particles comprising the active agent to be delivered to the cells are generated and delivered to the airway surfaces of the subject. Suitable vectors are typically viral vectors, including DNA viruses (wherein the nucleic acid active 25 agent is DNA) and RNA viruses, or retroviruses (wherein the nucleic acid active agent is RNA). Techniques for carrying out gene therapy are known and for the delivery of genetic materials to epithelial mucosa such as lung airway epithelial mucosa are known. See, e.g., F. Collins et al., U.S. Patent No. 5,240,846; I. Pastan, 5,166,059; R. Debs et Patent No. al., Application WO 93/12756; T. Friedmann, Science 244, 1275 (1989).

DNAs of the present invention, and oligonucleotides derived therefrom, are useful for screening for restriction fragment length polymorphism

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(RFLP) associated with disorders such as cystic fibrosis or other disorders potentially involving a defective P_{2U} receptor (or defective regulation thereof).

The present invention is explained in greater 5 detail in the following Examples. These Examples are for illustrative purposes only, and are not to be taken as limiting of the invention.

EXAMPLE 1

Expression of Human P20 Receptor in Human Astrocytoma Cells

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I. EXPERIMENTAL PROCEDURES

cDNA Cloning and Sequencing. Two degenerate oligonucleotide primers,

5'-AATGG(C/A/G)AC(C/T/A)TGGGA(G/A)GG(G/A)GA(C/T)GA(A/G)-3'. and

5'-GACGTG(C/G)AA(A/G)GGCAG(A/C)(A/C)AGC(A/T)GAGGGCGAA-3', from the N-terminal and transmembrane VI domains, respectively, of the murine P2U receptor sequence (Lustig et al., Proc. Natl. Acad. Sci. USA 90, 5113 (1993)), were used in a polymerase chain reaction to amplify products from a cDNA library constructed in lambda Uni-ZAP XR (Stratagene, La 20 Jolla, CA) from CF/T43 cell poly(A+) RNA. Amplification conditions were: 94°C, 1.5 min, 50°C, 1.5 min, 72°C, 2 min; 30 cycles. Products were cloned into pCR II (Invitrogen, San Diego, CA, USA) and screened by Southern blot using a random primer-labeled fragment corresponding 25 to bases 290--1097 (amino acid residues 8--276) of the murine P₂₀ sequence (probe P263).

A cloned PCR product of about 500 bases (probe D9) which hybridized with probe P263 was labeled by random priming and used to screen 7 x 10^5 recombinants of Hybridization was performed the CF/T43 cDNA library. QUIKHYB™ hybridization system (Stratagene) using the according to the manufacturer's instructions, except that 0.2x SSC/0.1% SDS was used for the high stringency wash. One plaque was identified which hybridized strongly with 35

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probe D9. After purification of phage by additional screening, pbluescript SK(-) was rescued by in vivo excision (Stratagene). An HT-29 cDNA library prepared in lambda gt10 was screened using probe P263, and insert from positive plaque-purified phage was subcloned into the Not I site of pbluescript. Plasmid DNA was purified by cesium chloride gradient centrifugation (Sambrook et al., Molecular Cloning: A Laboratory Manual (2d ed. 1989)) and both strands sequenced by dideoxy termination (Sequenase version 2.0; (United States Biochemical Corp., Cleveland, OH, USA).

Probe P263 was generated by high stringency PCR amplification of the murine P2U receptor clone, pP2R, obtained through the generosity of K. Lustig and D.

15 Julius (Univ. of California, San Francisco, CA, USA). The primers used to amplify P263 were:

5'-CTGGAATAGCACCATCAATGG-3' and 5'-GAGGTCAAGTGATCGGAAGG-3'.

Heterologous Expression. Α retroviral vector-containing plasmid, pLHP2USN, was constructed by 20 insertion of the cloned CF/T43 cDNA into the EcoRI and XhoI sites of pLXSN (A. Miller and G. Biotechniques 7, 980 (1989)). An amphotrophic packaging cell line, PA317, was used to produce the LHP2USN retroviral vector and a control vector containing Neor 25 only (LSN). Clonal human astrocytoma cells $1321N_1$, UNC strain, generously provided by T.K. Harden (UNC, Chapel Hill, NC), were plated at 150,000 cells per 60-mm dish and infected with LHP2USN or LSN (2 hr, Polybrene) and after 48 hr selected with 600 ug/ml G418 (Life Technologies, Inc., Gaithersburg, MD, USA). assay for receptor activity, intracellular calcium concentration was measured in confluent cells coverslips by fura-2 fluorescence microspectrofluorimetry as previously described (S. Mason et al., Br. J. Pharmacol. 103, 1649 (1991)). Measurement of cell inositol phosphate formation was as described by

H. Brown et al., Mol. Pharmacol. 40, 648 (1991), in accordance with known techniques

Northern and Southern Blots. Total cell and poly(A+) RNA were isolated by standard procedures or 5 purchased commercially (Clontech, Palo Alto, CA, USA). Human and mouse genomic DNA was extracted from cultured cells (Sambrook et al., supra) and digested (5 $U/\mu g$ DNA) with KpnI, EcoRI, Bam HI, or XbaI. RNA (formaldehyde denatured) or restriction fragments of genomic DNA were resolved on agarose gels, transferred to Nytran 10 nitrocellulose membranes (Schleicher & Schuell) described (Sambrook et al., supra), and UV-crosslinked (Stratagene). Prehybridization and hybridization blots with cDNA probes were performed using the QUIKHYB™ 15 hybridization system (Stratagene) as described above. Autoradiographs were done with Kodak XAR film and one intensifying screen at -80°C.

II. RESULTS AND DISCUSSION

A cDNA fragment from a human airway (CF/T43) cDNA library, amplified using degenerate primers based on 20 the murine P2U receptor sequence, was used to obtain a full-length plasmid clone, pHAP2U, from the same library, which contains the insert sequence given herein as SEQ ID Concomitantly, low stringency screening with the 25 murine P_{2U} partial cDNA, P263, of an HT-29 cDNA library produced a clone which, upon sequence analysis, was found to contain cDNA identical to the pHAP2U insert. deduced amino acid sequence of an open reading frame (SEQ ID NO: 1 and SEQ ID NO:2) present in the human cDNA bears substantial similarity (89% identity) to the murine P_{20} 30 receptor sequence reported by Lustig et al., supra, but similar considerably less to the chick purinoceptor (37% identity) described by T. Webb et al., FEBS Lett. 324, 219 (1993).

35 The pHAP2U insert (HP2U) sequence exhibits structural features typical of the family of G

protein-coupled receptors and common to both the chick and murine P_2 -receptors. These include: (1) hydrophobic domains, (2) consensus N-linked glycosylation sequences near the N-terminus, (3) a number of residues highly conserved among G protein-coupled receptors (e.g., Asn51, Asp79, Cys106 and Cys183), and (4) potential phosphorylation sites in the third intracellular and C-terminal domains. The putative P_{20} receptor sequences also have a potential site for phosphorylation by protein kinase C in the second intracellular loop. Alignment of 10 the human and murine P_{2U} receptor sequences reveals 4 additional residues, as well as a number of mismatched residues, in the C-terminal intracellular domain of the human clone. Like the murine and chick P2-receptors, the human P_{2U} receptor sequence does not appear to be closely related to any of the other cloned G protein-coupled receptors, including those for adenosine and cAMP.

A retrovirus expression system was employed to study the cloned CF/T43 HP2U cDNA in 1321N1 human 20 astrocytoma cells. The effect of extracellular nuceotides intracellular calcium, as measured by fluorescence, was assessed in CF/T43 cells, uninfected 1321N1 cells, and 1321N1 cells infected with retroviral vector containing either the cloned HP2U or no insert Exposure of CF/T43 cells to extracellular UTP 25 (10⁻⁴ M) resulted in an initial sharp increase in [Ca²⁺], which relaxed over 1-2 min to a prolonged plateau (data not shown). Application of UTP to CF/T43 cells in Ca₂₊-free medium induced a sharp increase in [Ca²⁺], which returned to baseline without an intervening plateau 30 phase. In contrast, extracellular UTP had no effect on uninfected 1321N1 cells (data not shown) or LSN-infected cells (data not shown), whereas carbachol elicited responses in each case. However, the [Ca2+], response of 1321N1 cells expressing the LHP2USN vector to UTP was 35 similar to that of CF/T43 cells in both Ca2.-free and Ca₂₊-containing media (data not shown). In

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experiment, the addition of ATP produced the same result observed with UTP. A response to extracellular nuceotides was seen in all trials with CF/T43 (peak [Ca²⁺]_i= 750 + 160 nM, n=10) and LHP2USN-expressing 1321N1 cells (peak [Ca²⁺]_i= 850 + 170 nM, n=6) but not with uninfected or LSN-infected controls.

To examine the pharmacological specifity of the expressed HP2U clone in 1321N1 cells, concentration-effect curves were generated 10 nucleotides and structural analogues active at Pox, Pox and P_{2U} receptors (data not shown). UTP and ATP equipotent, with EC₅₀ values of 1.4 and 1.0 uM. respectively, but UTP was slightly more efficacious. The P_{2X} agonist, 2MeSATP, and P_{2X} agonist, α, β -MeATP, had little effect. These concentration-effect relations are 15 very similar to those reported for heterologously expressed murine P2U (Lustig, et al., supra) and virtually indistinguishable from data (not shown) generated for the endogenous CF/T43 and HT-29 receptors. Clearly, the 20 agonist specifities fit the defininition of the P211 receptor.

Pretreatment of CF/T43 and HP2U-1321N1 cells with pertussis toxin inhibited ATP-stimulated intracellular Ca²⁺ mobilization in both cell-types by 20-30% (data not shown). These results are consistent with the effects of pertussis toxin on inositol phosphate accumulation induced by ATP or UTP in CF/T43 cells (Brown et al., supra (1991)), and on P_{2U} responses in other systems (Murphy and Tiffany, J. Biol. Chem. 265, 11615 30 (1990)).

The pertussis toxin sensitivity of HP2U-1321N1 cells, along with the observation that UTP/ATP raised $[Ca^{2+}]_i$ in the absence of extracellular Ca^{2+} , suggest that HP_{2U} is coupled to phospholipase C (PLC) via a G protein. To test this possibility more directly, we measured inositol phosphate formation. When these studies were first performed, very high levels of inositol phosphates

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were found to accumulate in HP_{20} -1321N1 cells which had not been exposed to exogenous nucleotides (Figure 1A). We hypothesized that during the lengthy labeling period 1321N1 cells released 5'-nucleotides into the medium in sufficient to "self-activate" the receptor. guantities The inclusion of apyrase in the medium during the baseline inositol cell-labeling step reduced the phosphate levels to values near those of controls (Figure Thus, it appears the cells are persistently stimulated by nucleotides in an autocrine fashion. 10 mode of release of nucleotides from the astrocytoma cells and the biological consequences of the persistent autocrine activation are unknown. To facilitate comparison with controls, the inositol phosphate studies performed on cells pretreated with 15 Consistent with coupling of the human P2v receptor to PLC, ATP and UTP increased inositol phosphate accumulation in 1321N1 cells expressing $LHP_{20}SN$ but not in uninfected or LSN-infected controls (data not shown).

P_{2U} receptor mRNA is widely distributed in human tissue: transcripts were detected in heart, liver, lung, and kidney, as reported for the mouse (Lustig et al., supra 1993), as well as placenta and skeletal muscle (data not shown). In addition, it was present in primary cultures of human respiratory epithelia and kidney proximal tubules, and in HSG-Pa salivary gland duct, CFT1 airway, and T84 colon cell lines.

were found to express more than one mRNA. Transcripts of 2.1 kb, 7.5 kb, and 9 kb were observed, and in some tissues all three RNAs were present. Although all of the bands in human tissues were found to cross hybridize with a full length murine P_{2U} receptor cDNA, none of the murine tissues analyzed by Lustig et al., supra, or mouse skeletal muscle contain more than one transcript. Human liver and other human cells, including primary cultures of proximal tubule epithelium, were found to express only

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the 2.1 kb message, but both CF/T43 and HT-29 cells express at least two transcripts.

Although Murphy and Tiffany, supra, have reported that HL60 cells may express more than one mRNA coding for functional P2U receptors, the significance of multiple transcripts is unknown. If all of the bands represent fully processed mRNAs they may be alternatively processed forms of the same gene or products of different genes. Alternative processing might produce transcripts that differ in their noncoding regions but code for On the other hand, since the identical proteins. structure of P2-receptor genes is unknown, the possibility remains that the different RNAs code for functionally different proteins, especially in light of the evidence presented below which suggests that at least two genes for P_{20} receptors exist in the human genome.

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Human genomic DNA was digested with restriction enzymes known not to cut in the cDNA fragments used as probes. Under stringent conditions, a human P20 receptor coding sequence probe, D9, recognized two discrete bands 20 in human genomic DNA cut with Kpn I, Eco RI, and Xba I, and three bands in DNA cut with Bam HI. The blot was then stripped and reprobed with a Kpn I fragment containing a segment of the 3' most bases of the HP_{2U} clone. If the bands are portions of the same gene, only 25 one would be associated with the 3' portion of the gene. In contrast, the HP20 Kpn I fragment recognized two bands in all of the digests (data not shown) suggesting that these bands are separate genes and not portions of the The downward shift of the more intensely 30 same gene. hybridizing band in the Kpn I digest is consistent with the presence of a Kpn I site between the sequences recognized by the two probes. The lack of a shift in the position of the less intensely hybridizing band of the Kpn I digest and the loss of one of three bands in the 35 Bam HI digest suggest that the second gene has lost the Kpn I site and that a Bam HI site has arisen in the

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portion recognized by the coding sequence probe. Both findings support the hypothesis that the probes are hybridizing to two unique genes.

A fragment from the coding region of the murine

5 P_{2U} receptor cDNA, probe P263, recognized a single band in
mouse genomic DNA digested with either Eco RI or Bam HI.
Whether these results correlate with the observation that
human tissues appear to express more than one species of
P_{2U} message, while only one P_{2U} receptor mRNA was found in
mouse tissues, remains to be resolved.

EXAMPLE 2

Expression of Murine P_{2U} Receptor in Human Leukemia Cells

I. EXPERIMENTAL PROCEDURES

15 Materials. pRc/CMV was purchased Invitrogen (San Diego, CA). K562 human leukemia cells (CCL-243) were obtained from American Type Culture Collection (Rockville, MD). Geneticin (G418) purchased from GIBCO (Grand Island, NY). Electroporation cuvettes (0.4 cm electrode gap) and the GENE PULSER™ obtained from electroporation chamber were Biorad (Richmond, CA). NYTRAN™ nylon membranes were purchased from Schleicher and Schuell (Keene, NJ). $[\alpha^{-32}P]$ ATP and $[\alpha^{-32}P]$ UTP were obtained from ICN Radiochemicals (Irvine, 25 CA). AVICEL™ microcrystalline silica gel thin layer chromatography plates containing a fluorescent indicator for nucleotides were obtained from Analtech (Newark, DE). N, N-dimethylformamide and 4-benzoylbenzoic acid were obtained from Aldrich (Milwaukee, WI). HIS-BIND™ resin was purchased from Novagen (Madison, WI). ATPγS was 30 purchased from Boehringer Mannheim (Indianapolis, IN) and 2-methylthioATP was purchased from Research Biochemicals Inc. (Natick, MA). Other reagents were purchased from Sigma (St. Louis, MO). Concentrations of nucleotide stock solutions were verified spectrophotometrically. 35

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Cell Culture. K562 cells were cultured in spinner flasks (58 rpm) in growth medium (pH 7.4) composed of RPMI-1640, 10% heat-inactivated fetal bovine serum, 2mM glutamine, 100 units/ml penicillin, and 100 μ g/ml streptomycin. The cell suspensions were maintained at 37°C in a humidified atmosphere of 5% CO₂ and 95% air at a density of 1x10⁵ to 1x10⁶ cells/ml.

Stable and Transient Transfections. To generate stable transfectants expressing the P20 receptor, K562 cells were transfected by electroporation 10 Toneguzzo et al., Mol. Cell. Biol. 6, 703 (1986)) with the neomycin-resistance plasmid pP2R1. pP2R1 is pRc/CMV containing a ~2.4 kb HindIII-NotI fragment of the P2U receptor cDNA, missing the first 6 nucleotides of the 5' untranslated region of the full length cDNA. cells were stably transfected with pRc/CMV. Cells were sedimented at 200 x g for 5 min at room temperature, rinsed once with phosphate-buffered saline (10 mM NaH2PO4, pH 7.4, 120 mM NaCl, 5 mM KCl), and resuspended in phosphate-buffered saline at a final concentration of 4 20 \times 10 7 cells/ml. The cells were chilled on ice for 10 min and then transferred in 0.5 ml aliquots to prechilled electroporation cuvettes. DNA (40 µg of pP2R1 linearized with PvuI) was added to the cell suspension, the cells 25 were subjected to a single electrical pulse (1 kV, 25 μ F) and the cuvettes were placed on ice. After 10 min, growth medium was added and the cells were incubated at 37°C. Fresh growth medium supplemented with 0.7 mg/ml G418 was added 48 h later, and the cells were incubated for 2 weeks to select for stable transfectants expressing 30 neomycin resistance. Clonal cell lines were then isolated by transferring single cells to individual wells of a 96-well culture plate: each well contained 0.2 ml of conditioned growth medium filtered from K562 stock cultures, 0.2 ml of fresh growth medium, and 0.4 mg/ml The clonal cell lines were cultured for three to four weeks in growth medium containing 0.4 mg/ml G418 and

assayed for P_{2U} receptor, the results shown are from one representative clonal cell line.

To generate transient transfectants, K562 cells were transfected as above with 40 μg unlinearized plasmid pCCM6xl or pP2R (Lustig et al., supra 1993). Cells were incubated for 48 h at 37°C in 100 mm culture dishes containing 12 ml of growth medium and assayed for P_{2U} receptor activity.

Cytoplasmic Free Calcium Measurements. Cells were sedimented at 200 x g for 5 min at room temperature 10 and washed once with 10 ml of Hepes-buffered saline (20 mM Hepes, pH 7.4, 120 mM NaCl, 5 mM KCl, 1 mM MgCl, and 1 mM CaCl2). Washed cells were suspended in Hepesbuffered saline at 5 x 105 cells/ml, and stored (for up to 4 h) at 37°C. The concentration of cytoplasmic free 15 [Ca²⁺];, measured was by dual-excitation calcium, spectrofluorometric analysis of cells loaded with fura-2, a fluorescent probe for calcium.

Northern analysis. Poly (A)* RNA was isolated from K562 cells with a FastTrack® mRNA isolation kit (Invitrogen) and electrophoresed in 0.8% agarose-6.5% formaldehyde gels. The mRNA was transferred by capillary flow to a nylon membrane, prehybridized for 15 min at 68°C in Quick-Hyb® (Stratagene), and hybridized for 1 h at 68°C with random-primer labeled DNA synthesized in vitro from the Hind III-Not I fragment of pP2R. The membrane was washed twice at room temperature in 2 X SSC containing 0.1% SDS, once at 60°C in 0.2 X SSC containing 0.1% SDS, and exposed to Kodak XAR film with an intensifying screen for 24 hr at -70°C.

Synthesis of $[\alpha^{-32}P]$ BzATP and $[\alpha^{-32}P]$ BZUTP. The following procedure was carried out in dim rom light to avoid photoactivation of the benzophenone moiety. 18 mg of 4-benzoylbenzoic acid (BzBz) and 42 mg of 1,1'-35 carbonyldimidazole were dissolved in 100 μ l of N,N-dimethylformamide and stirred in a tightly stoppered 5 ml polypropylene test tube for 15 min at room temperature.

Then, 0.5 ml of 3.3 μ M [α - 32 P]ATP (3000 Ci/mmol) was added slowly and the reaction mixture was stirred overnight (approx. 15 h) at room temperature in a loosely stopered test tube. The reaction mixture was centrifuged at 2000 5 x q for 5 min and the supernatant was added to a Sephadex LH-20 column (1.5 \times 45 cm) and eluted with 100 mM ammonium formate (pH 7.4) at a flow rate of 0.7 ml/min. The eluent was collected in 3.0 ml fractions and the fractions containing radioactivity were detected with a Geiger counter. Fractions 30-48 comprising the second 10 radioactive peak ($[\alpha^{-32}P]BzATP$) were pooled, lyophilized, reconstituted in 1 ml of H₂O and stored at -70°C. amount of radioactivity in the $[\alpha^{-32}P]$ BzATP sample was determined by liquid scintillation spectrometry and the 15 concentration of $[\alpha^{-32}P]$ BzATP and $[\alpha^{-32}P]$ ATP were equal. A 10-15% BzATP yield was obtained relative to the starting concentration of ATP. The purity of the $[\alpha$ -³²P]BzATP sample was assessed by thin layer chromatography (TLC) on silica gel plates containing a fluorescent indicator for nucleotides. A TLC plate was spotted with 1 μ l of the [α -³²P]BzATP sample and other lanes were spotted with 5 nanomoles of unlabeled ATP, BzATP (Sigma Chemical Co.) and BzBz. The plate was developed for 2 h temperature in a chromatography chamber room containing 100 ml of 1-butanol:H₂O:qlacial acetic acid (5:3:2, v/v/v) and the nonradioactive standards were visualized by exposing the plate to short wavelength ultraviolet light. The R_f values for ATP, BzATP and BzBz 0.98, respectively. were 0.65 and autoradiograph of the TLC plate indicated that the $[\alpha$ - 32 P]BzATP sample was >95% free of $[\alpha - ^{32}$ P]ATP contamination. Synthesis of $[\alpha^{-32}P]$ BzUTP was performed as for $[\alpha^{-32}P]$ BzATP except that 0.5 ml of 3.3 μ M [α - 32 P]UTP (3000 Ci/mmol) was substituted for $[\alpha^{-32}P]$ ATP in the reaction mixture. R, values of UTP and BzUTP were 0.21 and 0.70, 35 respectively.

5 GAGGATATCCTAATGATGATGATGATGTTGTTGTCTC (which contains an EcoRV site, stop codon, and the hexahistidine coding sequence) were incubated with 50 ng of pP2R, 20 nmoles of each dNTP, 1X reaction buffer (20 mM Tris-HCl, pH 8.2, 10 mM KCl, 6 mM $(NH_4)_2SO_4$, 1.5 mM MgCl₂, and 0.1% Triton X-100), and H₂O to bring the 10 reaction mixture to a 100 μ l volume. The reaction mixture was placed in a thermo-cycler and heated to 95°C for 5 min prior to the addition of 2.5 units of Pfu DNA polymerase (Stratagene), a thermostable enzyme that possesses both 5' to 3' polymerase and 3' 15 proofreading exonuclease activities. The mixture was overlayed with 50 μ l of mineral oil and subjected to 30 PCR cycles (95°C for 1.5 min, 55°C for 1.5 min, and 72°C for 2 min). The PCR product, a single 400 bp band, was subcloned into the unique BstXI-EcoRV 20 restriction site of pP2R to generate pP2HIS. K562 cells with stable P2HIS expression were produced by subcloning the HindIII-NotI restriction fragment of pP2HIS encoding the histidine-taged $P_{2\text{U}}$ receptor cDNA into pRC/CMV and transfecting K562 cells as described for pP2R1. 25 composition of pP2HIS was verified by DNA sequence analysis.

cells with stable P2HIS expression were purified according to procedure of Thom et al. *Biochem. J.* **168**, 187 (1977). The P_{2U} receptors were then photolabeled by incubating 1 mg of plasma membrane protein in 2 ml of Hepes-EDTA buffer (20 mM Hepes, pH 7.5, 1mM EDTA, 0.1 mM phenylmethylsulfonyl-fluoride, and 0.1 mM benzamidine) containing 5 nM [α - 32 P]BzATP or 5 nM [α - 32 P]BzUTP for 5 min at 37°C followed by irradiation with long wavelength UV light for 15 min at 4°C. The irradiated membranes were

washed twice with Hepes-EDTA buffer, pelleted centrifugation at 35,000 x g for 10 min at 4°C, and solubilized in 1 ml of urea buffer (6 M urea, 500 mM NaCl, 20 mM Tris-HCl, pH 7.9). The solubilized proteins 5 were applied to a Ni²⁺-charged Sepharose column (2.5 ml bed volume) and eluted with an imidazole gradient. Column eluents wre dialyzed against 10 mM Tris-HCl (pH 7.4), lyophilized, and resolubilized in 50 μ l of SDSsample buffer. The samples were electrophoresed on a 5-16% SDS-polyacrylamide gel for approximately 16 h at 5 10 milliamps. The gel was dried and exposed autoradiographic film for 1 to 3 days in order to visualize the radiolabeled proteins.

II. RESULTS AND DISCUSSION

ATP or UTP produced a two-fold increase in $[Ca^{2+}]_i$ in K562 cells stably or transiently transfected with P_{2U} receptor cDNA (data not shown). $[Ca^{2+}]_i$ was maximal within 5 s after nucleotide addition and then returned to the basal level within 1 min. Neither ATP nor UTP elevated $[Ca^{2+}]^i$ in untransfected K562 cells or in vector-transfected K562 cells. In transient transfectants, the largest P_{2U} receptor-mediated increases in $[Ca^{2+}]^i$ were observed 48 h after the transfection procedure.

In stable transfectants, the ATP- and UTP-25 mediated increases in [Ca2+] were not affected by depletion of extracellular Ca2+ by EGTA (data not shown), suggesting that P_{20} receptor activation causes the mobilization of intracellular calcium and not the influx of extracellular calcium. Consistent with this idea, ATP 30 and UTP also transiently reduced intracellular levels of phosphatidylinositol 4;5-bisphosphate (PIP2) in the stable transfectants by approximately 10%. Activation of endogenous P_{20} receptors in NG108-15 cells similarly leads 35 to the phospholipase C-dependent hydrolysis of PIP,, the generation of inositol 1,4,5-triphosphate (IP3), and the

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mobilization of calcium from intracellular stores (Lin et al., J. Neurochem. 60, 1115 (1993)). The relatively weak effect of ATP and UTP on PIP₂ hydrolysis and calcium mobilization in K562 cells transfected with P_{2U} receptor cDNA as compared to NG108-15 cells may reflect the low levels of IP₃ receptors in undifferentiated K562 cells.

It has been controversial whether ATP and UTP interact with the same receptor or whether the effect of each nucleotide is mediated by separate receptors. In stable transfectants, the EC_{50} values for ATP- and UTP-mediated increases in $[Ca^{2+}]_i$ were about 1.4 μ M and about 0.8 μ M, respectively (Figure 2). These responses are similar to those obtained in Xenopus oocytes expressing the P_{20} receptor where EC_{50} values of about 1 μ M were obtained for both nucleotides (Lustig et al., supra 1993). These findings demonstrate conclusively that the product of a single gene confers upon cells the ability to respond to both ATP and UTP.

K562 cells do not express an endogenous P_{20} receptor but do express a calcium-mobilizing P2r receptor 20 that is activated by ADP and inhibited by ATP. untransfected K562 cells, activation of the P_{2T} receptor. by ADP or 2-methylthioATP, but not 1 mM ATP, UTP, ATPYS, BzATP or BzUTP, elevated [Ca2+], (data not shown). ADP and 2-MeSATP do not appear to act as agonists of the P_{2U} 25 receptor, since the dose response curves for these virtually identical nucleotides were in transfectants and in vector-transfected K562 (Figure 2). Similarly, ADP and 2-methylthioATP did not stimulate 45Ca2+ efflux in oocytes expressing the cloned 30 receptor (Lustig et al., supra 1993). Hydrolysis of ATP may not be required for P2U receptor activation since the slowly-hydrolyzable ATP analog ATPγS was a relatively potent agonist (EC₅₀= about 9.6 μ M) of the P₂₀ receptor in 35 stable transfectants (Fig. 2) and in Xenopus oocytes (Lustig et al., supra 1993). Furthermore, β, γ methyleneATP, β , γ imidoATP, GTP, TTP, CTP, UDP, AMP and

adenosine, at concentrations as high as 1 mM, had no effect on $[Ca^{2+}]_i$; in untransfected K562 cells or stable transfectants, and did not prevent ATP-mediated increases in $[Ca^{2+}]_i$ in stable transfectants (not shown). Thus, these compounds do not appear to act as agonists or antagonists of the P_{2U} receptor.

The photoaffinity probes 3'-0-(4-benzoyl) benzoyl adenosine 5'-triphosphate (BzATP) and 3'-0-(4-benzoyl) benzoyl uridine 5'-triphosphate (BzUTP) also were agonists of the P_{2U} receptor (**Fig. 2**), although relatively weak agonists compared to ATP and UTP. BzATP has previously been used as an agonist and/or a photoaffinity probe for P_{2Z} receptors in transformed mouse fibroblasts, and mouse macrophages, P_{2Y} receptors in turkey erythrocytes, and P_{2U} receptors in bovine endothelial cells and NG108-15 cells. This, however, is the first report of BzUTP as a P_2 receptor agonist.

Isolation of the P_{2U} receptor and conclusive identification of its molecular weight were accomplished 20 with the aid of a hexahistidine tag attached to the carboxy terminus of the P_{20} receptor protein (P2HIS). Both ATP and UTP caused an increase in [Ca2+], and a decrease in PIP2 (not shown) in K562 cells that stably express P2HIS, indicating that the agonist specificity and signaling properties of the P_{2U} receptor were 25 unaltered by the hexahistidine tag. P2HIS was isolated by affinity chromatography on Ni2+-charged Sepharose columns. To increase detection of P2HIS, plasma membrane transfectants stable P2HIS proteins from photolabeled with $[\alpha^{-32}P]$ BzATP or $[\alpha^{-32}P]$ BzUTP. results indicated that a 53 kDa protein was selectively retained on the column until very high imidazole concentrations (200 mM) were used. The calculated molecular mass of the NG108-15 P_{20} receptor is 42 kDa, 35 suggesting that the mature receptor is glycosylated on one or both of two N-linked glycosyation consensus sites

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located in its putative amino-terminal extracellular domain.

The P_{2U} receptor cDNA encodes an mRNA of ~2.4kb in a variety of tissues and cell lines (Lustig et al, supra 1993). Northern blot analysis revealed that a P_{2U} receptor cDNA probe hybridized with an ~2.5 kb mRNA species in K562 cells that were transfected with P2R1 but did not hybridize with any transcript in untransfected K562 cells (data not shown). This suggests that the endogenous P_{2T} receptor in K562 cells is not significantly homologous to the P_{2U} receptor.

These results indicate that we have expressed a functional 53 kDa P_{2U} receptor in K562 human leukemia cells. In stable and transient transfectants, the P_{2U} receptor had the same agonist selectivity and signaling properties as in *Xenopus* oocytes and in NG108-15 cells, the source of the P_{2U} receptor cDNA. Thus, our findings define a mammalian model system for rapidly analyzing structure-function relationships, radioligand-binding and signal transduction properties of both wild-type and mutant P_{2U} receptors.

The foregoing examples are illustrative of the present invention, and are not to be construed as limiting thereof. The invention is defined by the following claims, with equivalents of the claims to be included therein.

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Boucher, Richard C. Weisman, Gary A. Turner, John T. Harden, Thomas K. Parr, Claude E. Sullivan, Daniel M. Erb, Laura Lustia, Kevin D.
- (ii) TITLE OF INVENTION: DNA Encoding the Human P2U Receptor and Null Cells Expressing the Same
- (iii) NUMBER OF SEQUENCES: 2
- (iv) CORRESPONDENCE ADDRESS:
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 - (F) ZIP: 28234
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk

 - (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0. Version #1.25
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE:
 - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:

 - (A) NAME: Sibley, Kenneth D.
 (B) REGISTRATION NUMBER: 31,665
 - (C) REFERENCE/DOCKET NUMBER: 5470-71
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 919-881-3140 (B) TELEFAX: 919-881-3175

 - (C) TELEX: 575102

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 1842 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single

-34-

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(ix) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 57..1181

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

(XT) SEQUENCE BESCHITTION. SEQ 15 No.1.							
GGAACCCGTG CAGGCGCTGA GCATCCTGAC CTGGAGAGCA GGGGCTGGTC AGGGCG							
CCC TGG AAT GAC ACC Pro Trp Asn Asp Thr 10	ATC AAT GGC ACC TGG Ile Asn Gly Thr Trp 15	104					
TAC AGG TGC CGC TTC Tyr Arg Cys Arg Phe 25	AAC GAG GAC TTC AAG Asn Glu Asp Phe Lys 30	152					
TCC TAC GGC GTG GTG Ser Tyr Gly Val Val 40	TGC GTG CTT GGG CTG Cys Val Leu Gly Leu 45	200					
		248					
TAT ATG TTC CAC CTG Tyr Met Phe His Leu 75	GCT GTG TCT GAT GCA Ala Val Ser Asp Ala 80	296					
CCG CTG CTG GTC TAT Pro Leu Leu Val Tyr 90	TAC TAC GCC CGC GGC Tyr Tyr Ala Arg Gly 95	344					
		392					
TGC AGC ATC CTC TTC Cys Ser Ile Leu Phe 120	CTC ACC TGC ATC AGC Leu Thr Cys Ile Ser 125	440					
GTC TTA CGA CCT CTG Val Leu Arg Pro Leu 135	CGC TCC CTG CGC TGG Arg Ser Leu Arg Trp 140	488					
CGC CGG GTG GCC GGG Arg Arg Val Ala Gly 155	GCC GTG TGG GTG TTG Ala Val Trp Val Leu 160	536					
A CF TT TS CL TT CP AT TC GV1 C	ATCCTGAC CTGGAGAGCA CCC TGG AAT GAC ACC Trp Asn Asp Thr 10 TAC AGG TGC CGC TTC Yr Arg Cys Arg Phe 25 TCC TAC GGC GTG GTG Tyr Gly Val Val 40 TTC TAC ATC TTC TTG Tyr Ile Phe Leu 55 TAT ATG TTC CAC CTG Yr Met Phe His Leu 75 TCG CTG CTG GTC TAT Tyr 90 ACG GTG CTC TGC AAG Thr Val Leu Val Tyr 90 ACG GTG CTC TGC AAG Thr Val Leu Cys Lys 105 TCC TAC AGC ATC CTC TTC TCC TTC TTC TTC TTC TTC TTC T	ATCCTGAC CTGGAGAGCA GGGGCTGGTC AGGGCG CCC TGG AAT GAC ACC ATC AAT GGC ACC TGG CPTO Trp Asn Asp Thr Ile Asn Gly Thr Trp 10 TAC AGG TGC CGC TTC AAC GAG GAC TTC AAG Cyr Arg Cys Arg Phe Asn Glu Asp Phe Lys 25 CCC TAC GGC GTG GTG TGC GTG CTT GGG CTG CPT Tyr Gly Val Val Cys Val Leu Gly Leu 40 CTC TAC ATC TTC TTG TGC CGC CTC AAG ACC Leu Tyr Ile Phe Leu Cys Arg Leu Lys Thr 55 AT ATG TTC CAC CTG GCT GTG TCT GAT GCA Cyr Met Phe His Leu Ala Val Ser Asp Ala 75 CCG CTG CTG GTC TAT TAC TAC GCC CGC GGC CPTO Leu Leu Val Tyr Tyr Tyr Ala Arg Gly 90 CCG GTG CTC TGC AAG CTG GTG CGC TTC CTC CHT Val Leu Cys Lys Leu Val Arg Phe Leu 105 CCC AGC ATC CTC TTC CTC ACC TGC ATC AGC Cys Ser Ile Leu Phe Leu Tys 120 CCC CGG GTG GCC GGG GCC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTA CGA CCT CTG CGC TCC CTG CGC TGG CTG TTG CTG CGC TGG CTG TTG CTG CGC CGG GTG GCC GGG GCC GTG TGG CTG TTG CTG CGC CGG GTG GCC GGG GCC GTG TGG CTG TTG CTG CGC CGG GTG GCC GGG GCC GTG TGG CTG TTG CTG CTG CTG CGC TGG CTG TTG CTG CTG CTG CGC TTG CTG CTG CTG CTG CTG					

GTG Val	CTG Leu	GCC Ala	TGC Cys	CAG Gln 165	. Ala	CCC Pro	GTG Val	CTC Leu	TAC Tyr 170	Phe	GTC Val	ACC Thr	ACC Thr	AGC Ser 175	GCG Ala	584
CGC Arg	GGG Gly	CCG Pro	CTA Leu 180	Thr	TGC Cys	CAC His	GAC Asp	ACC Thr 185	Ser	GCA Ala	CCC Pro	GAG Glu	CTC Leu 190	TTC Phe	AGC Ser	632
			Ala					Met	CTG Leu				Phe		GTG Val	680
CCC Pro	TTT Phe 210	Ala	GTC Val	ATC Ile	CTT Leu	GTC Val 215	TGT Cys	TAC Tyr	GTG Val	CTC Leu	ATG Met 220	GCT Ala	CGG Arg	CGA Arg	CTG Leu	728
									GGC Gly							776
									CTG Leu 250							824
TTC Phe	CTG Leu	CCA Pro	TTC Phe 260	His	GTC Val	ACC Thr	CGC Arg	ACC Thr 265	CTC Leu	TAC Tyr	TAC Tyr	TCC Ser	TTC Phe 270	CGC Arg	TCG Ser	872
CTG Leu	GAC Asp	CTC Leu 275	AGC Ser	TGC Cys	CAC His	ACC Thr	CTC Leu 280	AAC Asn	GCC Ala	ATC Ile	AAC Asn	ATG Met 285	GCC Ala	TAC Tyr	AAG Lys	920
GTT Val	ACC Thr 290	CGG Arg	CTG Leu	GCC Ala	AGT Ser	GCT Ala 295	AAC Asn	AGT Ser	TGC Cys	CTT Leu	GAC Asp 300	CCC Pro	GTG Val	CTC Leu	TAC Tyr	968
				Gln	Arg		Val	Arg	TTT Phe	Ala	Arg	Asp		Lys		1016
CCC Pro	ACT Thr	GGC Gly	Pro	AGC Ser 325	CCT Pro	GCC Ala	ACC Thr	CCG Pro	GCT Ala 330	CGC Arg	CGC Arg	ACG Thr	CTG Leu	GGC Gly 335	CTG Leu	1064
							Met		AGG Arg							1112
	Ser					Arg			TCC Ser		Pro					1160
			GAC Asp		Arg		TAGG	AGCA	GA A	CACT	TCAG	с ст	GTGC	AGGT		1211

TTATATTGGG AAGCTGTAGA GGACCAGGAC TTGTGCAGAC GCCACAGTCT CCCCAG	GATAT 1271
GGACCATCAG TGACTCATGC TGGATGACCC CATGCTCCGT CATTTGACAG GGGCTC	CAGGA 1331
TATTCACTCT GTGGTCCAGA GTCAACTGTT CCCATAACCC CTAGTCATCG TTTGTG	GTGTA 1391
TAAGTTGGGG GAATTAAGTT TCAAGAAAGG CAAGAGCTCA AGGTCAATGA CACCCC	CTGGC 1451
CTGACTCCCA TGCAAGTAGC TGGCTGTACT GCCAAGGTAC CTAGGTTGGA GTCCAG	GCCTA 1511
ATCAAGTCAA ATGGAGAAAC AGGCCCAGAG AGGAAGGTGG CTTACCAAGA TCACAT	TACCA 1571
GAGTCTGGAG CTGAGCTACC. TGGGGTGGGG GCCAAGTCAC AGGTTGGCCA GAAAAC	CCCTG 1631
GTAAGTAATG AGGGCTGAGT TTGCACAGTG GTCTGGAATG GACTGGGTGC CACGGT	TGGAC 1691
TTAGCTCTGA GGAGTACCCC CAGCCCAAGA GATGAACATC TGGGGACTAA TATCAA	ATAGA 1751
CCCATCTGGA GGCTCCCATG GGCTAGGAGC CAGTGTGAGG CTGTAACTTA TACTAA	AGGT 1811
TGTGTTGCCT GCTAAAAAAA AAAAAAAAAA A	1842

(2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 375 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Met Ala Ala Asp Leu Gly Pro Trp Asn Asp Thr Ile Asn Gly Thr Trp 1 5 10 15

Asp Gly Asp Glu Leu Gly Tyr Arg Cys Arg Phe Asn Glu Asp Phe Lys 20 25 30

Tyr Val Leu Leu Pro Val Ser Tyr Gly Val Val Cys Val Leu Gly Leu 35 40 45

Cys Leu Asn Ala Val Gly Leu Tyr Ile Phe Leu Cys Arg Leu Lys Thr $50 \hspace{1.5cm} 55 \hspace{1.5cm} 60$

Trp Asn Ala Ser Thr Thr Tyr Met Phe His Leu Ala Val Ser Asp Ala 65 70 75 80

Leu Tyr Ala Ala Ser Leu Pro Leu Leu Val Tyr Tyr Ala Arg Gly 85 90 95

Asp His Trp Pro Phe Ser Thr Val Leu Cys Lys Leu Val Arg Phe Leu 100 105 110

Phe Tyr Thr Asn Leu Tyr Cys Ser Ile Leu Phe Leu Thr Cys Ile Ser 115 Val His Arg Cys Leu Gly Val Leu Arg Pro Leu Arg Ser Leu Arg Trp Gly Arg Ala Arg Tyr Ala Arg Arg Val Ala Gly Ala Val Trp Val Leu Val Leu Ala Cys Gln Ala Pro Val Leu Tyr Phe Val Thr Thr Ser Ala Arg Gly Pro Leu Thr Cys His Asp Thr Ser Ala Pro Glu Leu Phe Ser Arg Phe Val Ala Tyr Ser Ser Val Met Leu Gly Leu Leu Phe Ala Val Pro Phe Ala Val Ile Leu Val Cys Tyr Val Leu Met Ala Arg Arg Leu 215 Leu Lys Pro Ala Tyr Gly Thr Ser Gly Gly Leu Pro Arg Ala Lys Arg Lys Ser Val Arg Thr Ile Ala Val Val Leu Ala Val Phe Ala Leu Cys Phe Leu Pro Phe His Val Thr Arg Thr Leu Tyr Tyr Ser Phe Arg Ser 265 Leu Asp Leu Ser Cys His Thr Leu Asn Ala Ile Asn Met Ala Tyr Lys 280 Val Thr Arg Leu Ala Ser Ala Asn Ser Cys Leu Asp Pro Val Leu Tyr 290 Phe Leu Ala Gly Gln Arg Leu Val Arg Phe Ala Arg Asp Ala Lys Pro Pro Thr Gly Pro Ser Pro Ala Thr Pro Ala Arg Arg Thr Leu Gly Leu Arg Arg Ser Asp Arg Thr Asp Met Gln Arg Ile Gly Asp Val Leu Gly Ser Ser Glu Asp Ser Arg Arg Thr Glu Ser Thr Pro Ala Gly Ser Glu Asn Thr Lys Asp Ile Arg Leu 370

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CLAIMS: .

- 1. Isolated DNA encoding an P_{2U} receptor selected from the group consisting of:
 - (a) isolated DNA consisting essentially of DNA which encodes the human P_{2U} receptor having the amino acid sequence given herein as SEQ ID NO:2;
 - (b) isolated DNA which hybridizes to isolated DNA of (a) above under conditions represented by a wash stringency of 0.3M NaCl, 0.03M sodium citrate, and 0.1% SDS at 60°C, and which encodes a human P_{2U} receptor; and
 - (c) isolated DNA differing from the isolated DNAs of (a) and (b) above in nucleotide sequence due to the degeneracy of the genetic code, and which encodes a human P_{2U} receptor.
- 2. Isolated DNA according to claim 1 consisting essentially of DNA which encodes the human P_{2U} receptor given herein as SEQ ID NO:2.
- 3. Isolated DNA according to claim 1 consisting 20 essentially of isolated DNA having the sequence given herein as SEQ ID NO:1.
 - 4. A recombinant DNA sequence comprising vector DNA and a DNA according to claim 1.
- 5. A recombinant DNA sequence according to claim 4, wherein said vector DNA comprises a vector selected from the group consisting of plasmids, adenoviruses, and cytomegaloviruses.
- 6. A recombinant DNA sequence according to claim 4, wherein said vector DNA comprises a baculovirus vector.

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- 7. A host cell containing a recombinant DNA sequence of claim 4 and capable of expressing the encoded protein.
- 8. A host cell according to claim 7, wherein said host cell is a mammalian cell.
 - 9. A host cell according to claim 7, wherein said host cell is selected from the group consisting of baby hamster kidney cells, mouse cells, human embryo cells, and chinese hamster ovary cells.
- 10. A transformed null cell which is essentially free of endogenous P_{2U} receptor expression, which null cell contains and expresses heterogenous DNA encoding an P_{2U} receptor, said heterogenous DNA selected from the group consisting of:
- (a) isolated DNA consisting essentially of DNA which encodes the human P_{2U} receptor having the amino acid sequence given herein as SEQ ID NO:2;
- (b) isolated DNA which hybridizes to isolated DNA of (a) above under conditions represented by a wash stringency of 0.3M NaCl, 0.03M sodium citrate, and 0.1% SDS at 60°C, and which encodes a P₂₀ receptor; and
- (c) isolated DNA differing from the isolated DNAs of (a) and (b) above in nucleotide sequence due to the degeneracy of the genetic code, and which encodes a P_{2U} receptor.
 - 11. A cell according to claim 10, which cell is essentially free of endogeneous P_{2x} and P_{2y} receptor expression.
- 12. A cell according to claim 10, which cell is 30 a human leukemia cell.

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- 13. A cell according to claim 10, which cell is a K562 human leukemia cell.
- 14. A cell according to claim 10, which cell is a human astrocytoma cell.
- 5 15. A cell according to claim 10, which cell is a $1321N_1$ human astrocytoma cell.
 - 16. A method of detecting compounds which bind to the $P_{2\text{U}}$ receptor, comprising:
- (a) providing a cell which contains and expresses heterogenous DNA encoding a heterogenous P_{2U} receptor, which cell secretes endogenous ATP in response to stimulation of said P_{2U} receptor, said heterogenous DNA selected from the group consisting of:
 - (i) isolated DNA consisting essentially of DNA which encodes the human P_{2U} receptor having the amino acid sequence given herein as SEQ ID NO:2;
 - (ii) isolated DNA which hybridizes to isolated DNA of (i) above under conditions represented by a wash stringency of 0.3M NaCl, 0.03M sodium citrate, and 0.1% SDS at 60°C, and which encodes a P_{2U} receptor; and
 - (iii) isolated DNA differing from the isolated DNAs of (i) and (ii) above in nucleotide sequence due to the degeneracy of the genetic code, and which encodes a P_{2U} receptor;
 - (b) removing essentially all endogenous ATP secreted by said cell from said cell;
 - (c) contacting a compound to said cell; and then
- 30 (d) detecting the binding of said compound to said P_{2U} receptor.

- 17. A method according to claim 16, wherein said removing step is carried out by enzymatically degrading essentially all of said endogenous ATP.
- 18. A method according to claim 16, wherein 5 said removing step is carried out by separating said cell from essentially all of said endogenous ATP.
- 19. A method according to claim 16, wherein said contacting step is carried out in an aqueous solution, and wherein said removing step is carried out by including hexokinase and glucose in said aqueous solution in an amount sufficient so that essentially all of said ATP is utilized to phosphorylate said glucose.
 - 20. A method according to claim 16, wherein said contacting step is carried out in an aqueous solution, and wherein said removing step is carried out by including apyrase in said aqueous solution in an amount effective to degrade said ATP.
 - 21. A method according to claim 16, wherein said removing step is carried out by washing said cell.
- 20 22. A method of enhancing cell growth, comprising,

transforming a cell with a vector capable of expressing a P_{2U} receptor in said cell in an amount sufficient to enhance cell growth by autostimulation of said P_{2U} receptor, wherein said cell secretes ATP, and wherein cell growth is enhanced by stimulation of said P_{2U} receptor.

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23. A method according to claim 22, wherein said cell is a mammalian cell.

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24. A method according to claim 22, further comprising the step of growing said cell in culture.

- 25. A method according to claim 22, wherein said vector contains and expresses DNA encoding a P_{2U} receptor selected from the group consisting of:
- (a) isolated DNA consisting essentially of DNA which encodes the human P_{2U} receptor having the amino acid sequence given herein as SEQ ID NO:2;
- (b) isolated DNA which hybridizes to isolated 10 DNA of (a) above under conditions represented by a wash stringency of 0.3M NaCl, 0.03M sodium citrate, and 0.1% SDS at 60°C, and which encodes a P_{2U} receptor; and
- (c) isolated DNA differing from the isolated DNAs of (a) and (b) above in nucleotide sequence due to the degeneracy of the genetic code, and which encodes a P_{2U} receptor.
 - 26. A method of inhibiting cell growth, comprising,

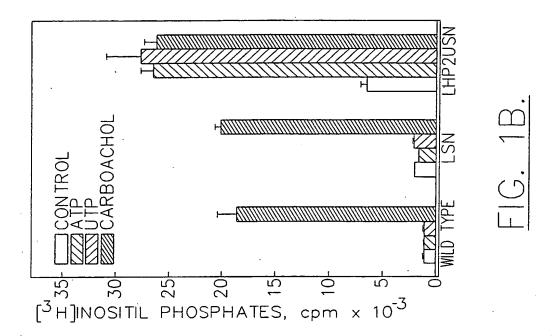
transforming a cell with a vector capable of 20 expressing a P_{2U} receptor in said cell in an amount sufficient to inhibit cell growth by autostimulation of said P_{2U} receptor, wherein said cell secretes ATP, and wherein cell growth is inhibited by stimulation of said P_{2U} receptor.

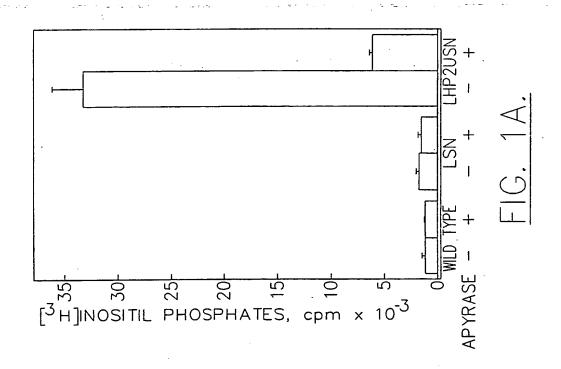
- 25 27. A method according to claim 26, wherein said cell is a mammalian cell.
 - 28. A method according to claim 26, wherein said vector contains and expresses DNA encoding a P_{2U} receptor selected from the group consisting of:
- 30 (a) isolated DNA consisting essentially of DNA which encodes the human P_{2U} receptor having the amino acid sequence given herein as SEQ ID NO:2;

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- (b) isolated DNA which hybridizes to isolated DNA of (a) above under conditions represented by a wash stringency of 0.3M NaCl, 0.03M sodium citrate, and 0.1% SDS at 60°C, and which encodes a P_{2U} receptor; and
- (c) isolated DNA differing from the isolated DNAs of (a) and (b) above in nucleotide sequence due to the degeneracy of the genetic code, and which encodes a $P_{2\text{U}}$ receptor.

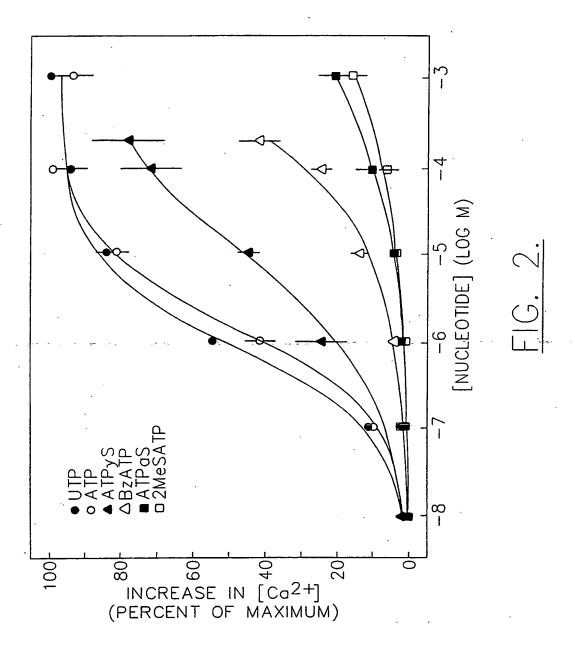




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B. FIEI	LDS SEARCHED		
	locumentation searched (classification system followe 435/6, 7.1, 69.1; 536/23.1	d by classification symbols)	
Documental	tion searched other than minimum documentation to th	e extent that such documents are i	ncluded in the fields searched
APS, Me	data base consulted during the international search (needline, Biosis, WPI erms: P2U receptor, ATP receptor, ligand, cell		cticable, search terms used)
C. DOC	CUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where a	ppropriate, of the relevant passag	Relevant to claim No.
Υ	Molecular Pharmacology, Volume al, "Evidence that UTP and ATP through a Common Extracellular Human Airway Epithelial Cells", p 651.	Regulate Phospholipa: 5'-Nucleotide Recepto	se C or in
X Y	Proceedings of the National Acad 90, Number 11, issued 01 Ju "Expression Cloning of an ATI Neuroblastoma Cells", pages 511 5116.	ine 1993, Lustig et P Receptor from Mo	al, 2, buse 3, 12-28
X Furth	er documents are listed in the continuation of Box C	. See patent family an	nnex.
"A" doc to b "E" earl "L" doc cite spec "O" doc mee		"Y" document of particular rele considered novel or cannot b when the document is taken "Y" document of particular rele considered to involve an combined with one or more being obvious to a person sk	evance; the claimed invention cannot be the considered to involve an inventive step alone evance; the claimed invention cannot be inventive step when the document is other such documents, such combination cilled in the art
the	rument published prior to the international filing date but later than priority date claumed	*&* document member of the sar	
	MBER 1994	Date of mailing of the internation 18 JAN1	
Commission Box PCT	nailing address of the ISA/US ner of Patents and Trademarks	Authorized officer Sally P. Teng	0. Kin 3a 1

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(703) 308-0196

Facsimile No.

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.		
Y	Trends in Pharmacological Sciences, Volume 12, Number 4, issued April 1991, O'Connor et al, "Further Subclassification of ATP Receptors Based on Agonist Studies", pages 137-141, see pages 138 and 139.	1-28		
Y	Federation of European Biochemical Societies, Volume 324, Number 2, issued June 1993, Webb et al., "Cloning and Functional Expression of a Brain G Protein-Coupled ATP Receptor", pages 219-225, see pages 220-223.			
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